GenCore version 5.1.7 Copyright (c) 1993 - 2006 Biocceleration Ltd.

OM nucleic - nucleic search, using sw model

Run on: April 14, 2006, 16:49:04; Search time 363 Seconds

(without alignments)

9470.535 Million cell updates/sec

Title: US-10-086-623-5

Perfect score: 1934

Sequence: 1 ttgtaccgaagagatgagac.....atcgacgtaactggaaaccg 1934

Scoring table: IDENTITY_NUC

Gapop 10.0 , Gapext 1.0

Searched: 1303057 seqs, 888780828 residues

Total number of hits satisfying chosen parameters: 2606114

Minimum DB seq length: 0

Maximum DB seq length: 200000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Database : Issued_Patents_NA:*

- 1: /cgn2_6/ptodata/1/ina/1_COMB.seq:*
- 2: /cgn2_6/ptodata/1/ina/5_COMB.seq:*
- 3: /cgn2 6/ptodata/1/ina/6A_COMB.seq:*
- 4: /cgn2 6/ptodata/1/ina/6B_COMB.seq:*
- 5: /cgn2_6/ptodata/1/ina/H_COMB.seq:*
- 6: /cgn2_6/ptodata/1/ina/PCTUS_COMB.seq:*
- 7: /cgn2_6/ptodata/1/ina/PP_COMB.seq:*
- 8: /cgn2_6/ptodata/1/ina/RE_COMB.seq:*
- 9: /cgn2_6/ptodata/1/ina/backfiles1.seq:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		8				
Result		Query				
No.	Score	Match	Length	DB	ID	Description
1	1934	100.0	1934	-	US-09-438-046-5	Sequence 5, Appli
2	1934	100.0	2253	3	US-09-438-046-7	Sequence 7, Appli
3	1491.8	77.1	1882	3	US-09-457-066-36	Sequence 36, Appl
4	1491.8	77.1	1882	3	US-09-540-224-1	Sequence 1, Appli
5	1491.8	77.1	1882	3	US-09-564-595D-1	Sequence 1, Appli
6	1491.8	77.1	1882	3	US-09-706-968-36	Sequence 36, Appl
7	1491.8	77.1	1882	3	US-09-808-972-1	Sequence 1, Appli
8	1491.8	77.1	1882	3	US-10-139-583-36	Sequence 36, Appl

```
1491.8
                 77.1
                        1882
                                                               Sequence 1, Appli
                               3
                                  US-10-039-847A-1
   10
       1491.8
                 77.1
                        1882
                                  US-10-226-559-1
                                                               Sequence 1, Appli
                               3
   11 . 1491.8
                 77.1
                        1882
                                                               Sequence 36, Appl
                                  US-09-541-752-36
       1491.8
                 77.1
                        1882
   12
                                  US-09-876-813A-1
                                                               Sequence 1, Appli
                               3
        795.4
                 41.1
                        1472
                                  US-09-540-224-3
   13
                               3
                                                               Sequence 3, Appli
        795.4
                 41.1
                        1472
   14
                               3
                                  US-09-564-595D-52
                                                               Sequence 52, Appl
   15
        795.4
                        1472
                                  US-09-808-972-3
                                                               Sequence 3, Appli
                 41.1
   16
        795.4
                        1472
                 41.1
                                  US-10-039-847A-3
                                                               Sequence 3, Appli
        795.4
                        1472
                                                               Sequence 3, Appli
   17
                 41.1
                                  US-10-226-559-3
                               3
   18
        795.4
                 41.1
                        1472
                                  US-09-876-813A-52
                               3
                                                               Sequence 52, Appl
   19
                        1110
                                  US-09-564-595D-6
        697.6
                 36.1
                               3
                                                               Sequence 6, Appli
   20
        697.6
                 36.1
                        1110
                                                               Sequence 6, Appli
                                  US-09-876-813A-6
                               3
   21
          662
                 34.2
                          690
                               3
                                  US-09-438-046-3
                                                               Sequence 3, Appli
                                                               Sequence 1, Appli
        355.2
                 18.4
                          360
   22
                                  US-09-438-046-1
                               3
   23
        213.8
                 11.1
                          256
                                  US-09-222-575-13
                                                               Sequence 13, Appl
                               3
        213.8
                 11.1
                          256
   24
                               3
                                  US-09-389-681-13
                                                               Sequence 13, Appl
C
        213.8
                 11.1
                         256
                                                               Sequence 13, Appl
   25
                               3
                                  US-09-620-405B-13
C
        213.8
                 11.1
                          256
                                  US-09-339-338-13
   26
                               3
                                                               Sequence 13, Appl
C
        213.8
                 11.1
                                  US-09-433-826B-13
   27
                          256
                                                               Sequence 13, Appl
C
   28
        213.8
                 11.1
                         256
                                  US-09-604-287A-13
                                                               Sequence 13, Appl
C
                         256
        213.8
                 11.1
                                  US-09-285-480-13
   29
                               3
                                                               Sequence 13, Appl
C
        213.8
                 11.1
                         256
                                  US-09-834-759-13
   30
                               3
                                                               Sequence 13, Appl
C
                          256
        213.8
                 11.1
                                  US-09-590-751A-13
   31
                                                               Sequence 13, Appl
   32
                 11.1
                         256
        213.8
                                  US-09-551-621-13
                                                               Sequence 13, Appl
                               3
C
                         256
   33
        213.8
                 11.1
                               3
                                  US-09-551-621A-13
                                                               Sequence 13, Appl
C
   34
                          256
                                                               Sequence 13, Appl
        213.8
                 11.1
                               3
                                  US-10-076-622-13
                                                               Sequence 6, Appli
                  9.4
                        1035
   35
        181.8
                               3
                                  US-09-457-066-6
                                                               Sequence 6, Appli
        181.8
                        1035
   36
                               3
                  9.4
                                  US-09-706-968-6
        181.8
                  9.4
                        1035
                                  US-10-139-583-6
                                                               Sequence 6, Appli
   37
                               3
        181.8
                  9.4
                        1035
                                                               Sequence 6, Appli
   38
                               3
                                  US-09-541-752-6
        123.2
                         3571
                                                               Sequence 42, Appl
   39
                  6.4
                                  US-09-457-066-42
                         3571
   40
        123.2
                  6.4
                                  US-09-564-595D-34
                                                               Sequence 34, Appl
        123.2
                  6.4
                        3571
                                  US-09-706-968-42
                                                               Sequence 42, Appl
   41
                               3
   42
        123.2
                        3571
                                                               Sequence 3, Appli
                  6.4
                               3
                                  US-09-823-033-3
                         3571
                                                               Sequence 42, Appl
        123.2
                                  US-10-139-583-42
   43
                  6.4
   44
        123.2
                  6.4
                         3571
                               3
                                                               Sequence 42, Appl
                                  US-09-541-752-42
                                                               Sequence 3, Appli
                         3571
   45
        123.2
                               3
                                  US-09-695-121-3
                  6.4
```

ALIGNMENTS

```
RESULT 1
US-09-438-046-5
; Sequence 5, Application US/09438046
; Patent No. 6706687
; GENERAL INFORMATION:
; APPLICANT: ERIKSSON, Ulf
; APPLICANT: AASE, Karin
; APPLICANT: LEE, Xuri
; APPLICANT: PONTN, Annica
; APPLICANT: UUTELA, Marko
; APPLICANT: ALITALO, Kari
; APPLICANT: OESTMAN, Arne
; APPLICANT: HELDIN, Carl-Henrik
; TITLE OF INVENTION: PLATELET-DERIVED GROWTH FACTOR D, DNA CODING
; TITLE OF INVENTION: THEREFOR, AND USES THEREOF
```

GenCore version 5.1.7 Copyright (c) 1993 - 2006 Biocceleration Ltd.

OM nucleic - protein search, using frame_plus_n2p model

Ruń on: April 14, 2006, 14:31:10; Search time 12.9 Seconds

(without alignments)

2478.989 Million cell updates/sec

Title: US-10-086-623-5

Perfect score: 3446

Sequence: 1 ttgtaccgaagagatgagac.....atcgacgtaactggaaaccg 1934

Scoring table: BLOSUM62

Xgapop 10.0 , Xgapext 0.5
Ygapop 10.0 , Ygapext 0.5
Fgapop 6.0 , Fgapext 7.0
Delop 6.0 , Delext 7.0

Searched: 572060 seqs, 82675679 residues

Total number of hits satisfying chosen parameters: 1144120

Minimum DB seq length: 0

Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Maximum Match 100%

Listing first 45 summaries

Command line parameters:

- -MODEL=frame+ n2p.model -DEV=xlp
- -Q=/abss/ABSSWEB_spool/US10086623/runat_14042006_123917_255/app_query.fasta_1
- -DB=Issued Patents AA -QFMT=fastan -SUFFIX=rai -MINMATCH=0.1 -LOOPCL=0
- -LOOPEXT=0 -UNITS=bits -START=1 -END=-1 -MATRIX=blosum62 -TRANS=human40.cdi
- -LIST=45 -DOCALIGN=200 -THR_SCORE=pct -THR_MAX=100 -THR_MIN=0 -ALIGN=15
- -MODE=LOCAL -OUTFMT=pto -NORM=ext -HEAPSIZE=500 -MINLEN=0 -MAXLEN=200000000
- -HOST=abss02p -USER=US10086623_@CGN_1_1_71_@runat_14042006_123917_255 -NCPU=6
- -ICPU=3 -NO_MMAP -NEG_SCORES=0 -WAIT -DSPBLOCK=100 -LONGLOG -DEV_TIMEOUT=120
- -WARN TIMEOUT=30 -THREADS=1 -XGAPOP=10 -XGAPEXT=0.5 -FGAPOP=6 -FGAPEXT=7
- -YGAPOP=10 -YGAPEXT=0.5 -DELOP=6 -DELEXT=7

Database : Issued Patents AA:*

- 1: /cgn2_6/ptodata/1/iaa/5_COMB.pep:*
- 2: /cgn2_6/ptodata/1/iaa/6_COMB.pep:*
- 3: /cgn2_6/ptodata/1/iaa/H_COMB.pep:*
- 4: /cgn2_6/ptodata/1/iaa/PCTUS_COMB.pep:*
- 5: /cgn2_6/ptodata/1/iaa/RE_COMB.pep:*
- 6: /cgn2_6/ptodata/1/iaa/backfiles1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

		8					
Result		Query					
No.	Score	Match	Length	DB	ID	Description	
1	1742	50.6	322	2	US-09-438-046-6	Sequence 6, Appli	
2	1742	50.6	370	2	US-09-457-066-37	Sequence 37, Appl	
3	1742	50.6	370	2	US-09-540-224-2	Sequence 2, Appli	
4	1742	50.6	370	2	US-09-564-595D-2	Sequence 2, Appli	
5	1742	50.6	370	2	US-09-706-968-37	Sequence 37, Appl	
6	1742	50.6	370	2	US-09-808-972-2	Sequence 2, Appli	
7	1742	50.6	370	2	US-09-823-033-5	Sequence 5, Appli	
8	1742	50.6	370	2	US-09-438-046-8	Sequence 8, Appli	
9	1742	50.6	370	2	US-10-139-583-37	Sequence 37, Appl	
10	1742	50.6	370	2	US-10-039-847A-2	Sequence 2, Appli	
11	1742	50.6	370	2	US-10-226-559-2	Sequence 2, Appli	
12	1742	50.6	370	2	US-09-541-752-37	Sequence 37, Appl	
13	1742	50.6	370	2	US-09-695-121-5	Sequence 5, Appli	
14	1742	50.6	370	2	US-09-876-813A-2	Sequence 2, Appli	
15	1561	45.3	370	2	US-09-540-224-4	Sequence 4, Appli	
16	1561	45.3	370	2	US-09-564-595D-53	Sequence 53, Appl	
17	1561	45.3	370	2	US-09-808-972-4	Sequence 4, Appli	
18	1561	45.3	370	2	US-10-039-847A-4	Sequence 4, Appli	
19	1561	45.3	370	2	US-10-226-559-4	Sequence 4, Appli	
20	1561	45.3	370	2	US-09-876-813A-53	Sequence 53, Appl	
21	1403	40.7	317	2	US-09-564-595D-56	Sequence 56, Appl	
22	1357.5	39.4	316	2	US-09-564-595D-55	Sequence 55, Appl	
23	1095	31.8	303	2	US-09-564-595D-57	Sequence 57, Appl	
24	1061.5	30.8	302	2	US-09-564-595D-54	Sequence 54, Appl	
25	1037	30.1	200	2	US-09-438-046-4	Sequence 4, Appli	
26	750.5	21.8	345	2	US-09-457-066-43	Sequence 43, Appl	
27	750.5	21.8	345	2	US-09-564-595D-35	Sequence 35, Appl	
28	750.5	21.8	345	2	US-09-706-968-43	Sequence 43, Appl	
29	750.5	21.8	345	2	US-09-823-033-4	Sequence 4, Appli	
30	750.5	21.8	345	2	US-10-139-583-43	Sequence 43, Appl	
31	750.5	21.8	345	2	US-09-541-752-43	Sequence 43, Appl	
32	750.5	21.8	345	2	US-09-695-121-4	Sequence 4, Appli	
33	750.5	21.8	345	2	US-09-876-813A-35	Sequence 35, Appl	
34	741	21.5	374	2	US-09-468-647A-118	Sequence 118, App	
35	739.5	21.5	323	2	US-09-468-647A-1	Sequence 1, Appli	
36	739.5	21.5	345	2	US-09-040-220D-2	Sequence 2, Appli	
37	739.5	21.5	345	2	US-09-457-066-2	Sequence 2, Appli	
38	739.5	21.5	345	2	US-09-265-686-2	Sequence 2, Appli	
39	739.5	21.5	345	2	US-09-540-224-5	Sequence 5, Appli	
40	739.5	21.5	345	2	US-09-564-595D-33	Sequence 33, Appl	
41	739.5	21.5	345	2	US-09-706-968-2	Sequence 2, Appli	
42	739.5	21.5	345	2	US-09-723-749-2	Sequence 2, Appli	
43	739.5	21.5	345	2	US-09-823-033-2	Sequence 2, Appli	
		0.5	245	_	TTO AO 460 6473 3	Compande 2 Annli	

ALIGNMENTS

345 2 US-09-468-647A-2

345 2 US-09-468-647A-101

Sequence 2, Appli

Sequence 101, App

RESULT 1 US-09-438-046-6

44

45

21.5

739.5

739.5 21.5

[;] Sequence 6, Application US/09438046

[;] Patent No. 6706687

```
; GENERAL INFORMATION:
 ; APPLICANT: ERIKSSON, Ulf
 ; APPLICANT: AASE, Karin
 ; APPLICANT: LEE, Xuri
 ; APPLICANT: PONTN, Annica
 ; APPLICANT: UUTELA, Marko
 ; APPLICANT: ALITALO, Kari
    APPLICANT: OESTMAN, Arne
 ; APPLICANT: HELDIN, Carl-Henrik
    TITLE OF INVENTION: PLATELET-DERIVED GROWTH FACTOR D, DNA CODING
    TITLE OF INVENTION: THEREFOR, AND USES THEREOF
  FILE REFERENCE: Ulf Eriksson et al
 ; CURRENT APPLICATION NUMBER: US/09/438,046
 ; CURRENT FILING DATE: 1999-11-10
 ; EARLIER APPLICATION NUMBER: 60/107,852
    EARLIER FILING DATE: 1998-11-10
 ; EARLIER APPLICATION NUMBER: 60/113,997
 ; EARLIER FILING DATE: 1999-12-28
 ; EARLIER APPLICATION NUMBER: 60/150,604
    EARLIER FILING DATE: 1999-08-26
 ; EARLIER APPLICATION NUMBER: 60/157,108
 ; EARLIER FILING DATE: 1999-10-04
 ; EARLIER APPLICATION NUMBER: 60/157,756
 ; EARLIER FILING DATE: 1999-10-05
; NUMBER OF SEQ ID NOS: 31
 ; SOFTWARE: PatentIn Ver. 2.0
 ; SEQ ID NO 6
 ; LENGTH: 322
     TYPE: PRT
     ORGANISM: Homo sapiens
 US-09-438-046-6
 Alignment Scores:
                                    Length:
                                                 322
 Pred. No.:
                      4.43e-191
                                                 322
                                    Matches:
                      1742.00
 Score:
 Percent Similarity:
                      100.0%
                                    Conservative:
 Best Local Similarity: 100.0%
                                    Mismatches:
                                    Indels:
 Query Match:
                       50.6%
                                    Gaps:
 DB:
 US-10-086-623-5 (1-1934) x US-09-438-046-6 (1-322)
            1 TTGTACCGAAGAGATGAGACCATCCAGGTGAAAGGAAACGGCTACGTGCAGAGTCCTAGA 60
 Qу
              1 LeuTyrArgArgAspGluThrIleGlnValLysGlyAsnGlyTyrValGlnSerProArg 20
 Db
           61 TTCCCGAACAGCTACCCCAGGAACCTGCTCCTGACATGGCGGCTTCACTCTCAGGAGAAT 120
 Qу
              21 PheProAsnSerTyrProArgAsnLeuLeuLeuThrTrpArgLeuHisSerGlnGluAsn 40
 Db
          121 ACACGGATACAGCTAGTGTTTGACAATCAGTTTGGATTAGAGGAAGCAGAAAATGATATC 180
 Qy
              41 ThrArgIleGlnLeuValPheAspAsnGlnPheGlyLeuGluGluAlaGluAsnAspIle 60
 Db
          181 TGTAGGTATGATTTTGTGGAAGTTGAAGATATATCCGAAACCAGTACCATTATTAGAGGA 240
 Qy
              61 CysArgTyrAspPheValGluValGluAspIleSerGluThrSerThrIleIleArgGly 80
 Db
```

Qy	241		300
Db	81	ArgTrpCysGlyHisLysGluValProProArgIleLysSerArgThrAsnGlnIleLys	100
Qy	301	ATCACATTCAAGTCCGATGACTACTTTGTGGCTAAACCTGGATTCAAGATTTATTCT	360
Db	101	IleThrPheLysSerAspAspTyrPheValAlaLysProGlyPheLysIleTyrTyrSer	120
Qy	361	TTGCTGGAAGATTTCCAACCCGCAGCAGCTTCAGAGACCAACTGGGAATCTGTCACAAGC	420
Db	121	LeuLeuGluAspPheGlnProAlaAlaAlaSerGluThrAsnTrpGluSerValThrSer	140
Qy	421	TCTATTTCAGGGGTATCCTATAACTCTCCATCAGTAACGGATCCCACTCTGATTGCGGAT	480
Db	141	SerIleSerGlyValSerTyrAsnSerProSerValThrAspProThrLeuIleAlaAsp	160
Qу	481	GCTCTGGACAAAAAATTGCAGAATTTGATACAGTGGAAGATCTGCTCAAGTACTTCAAT	540
Db	161	AlaLeuAspLysLysIleAlaGluPheAspThrValGluAspLeuLeuLysTyrPheAsn	180
QУ	541	CCAGAGTCATGGCAAGAAGATCTTGAGAATATGTATCTGGACACCCCTCGGTATCGAGGC	600
Db	181	ProGluSerTrpGlnGluAspLeuGluAsnMetTyrLeuAspThrProArgTyrArgGly	200
Qy	601	AGGTCATACCATGACCGGAAGTCAAAAGTTGACCTGGATAGGCTCAATGATGATGCCAAG	660
Db	201	ArgSerTyrHisAspArgLysSerLysValAspLeuAspArgLeuAsnAspAspAlaLys	220
Qу	661	CGTTACAGTTGCACTCCCAGGAATTACTCGGTCAATATAAGAGAAGAGCTGAAGTTGGCC	720
Db	221	ArgTyrSerCysThrProArgAsnTyrSerValAsnIleArgGluGluLeuLysLeuAla	240
Qу	721	AATGTGGTCTTCTTTCCACGTTGCCTCCTCGTGCAGCGCTGTGGAGGAAATTGTGGCTGT	780
Db	241	AsnValValPhePheProArgCysLeuLeuValGlnArgCysGlyGlyAsnCysGlyCys	260
Qу	781	GGAACTGTCAACTGGAGGTCCTGCACATGCAATTCAGGGAAAACCGTGAAAAAGTATCAT	840
Db	261	GlyThrValAsnTrpArgSerCysThrCysAsnSerGlyLysThrValLysLysTyrHis	280
Qу	841	GAGGTATTACAGTTTGAGCCTGGCCACATCAAGAGGGGGGGTAGAGCTAAGACCATGGCT	900
Db	281	GluValLeuGlnPheGluProGlyHisIleLysArgArgGlyArgAlaLysThrMetAla	300
Qy	901	CTAGTTGACATCCAGTTGGATCACCATGAACGATGCGATTGTATCTGCAGCTCAAGACCA	960
Db	301	LeuValAspIleGlnLeuAspHisHisGluArgCysAspCysIleCysSerSerArgPro	320
Qу	961	CCTCGA 966	
Db	321	ProArg 322	

RESULT 2

US-09-457-066-37

[;] Sequence 37, Application US/09457066 ; Patent No. 6432673